DOUBLE STRANDED RNA

Amendments to the Claims

This listing of claims replaces all prior versions, and listings, of claims in the aboveidentified application:

- 1. (currently amended) A method for attenuating the expression of a target gene in a vertebrate cell *in vivo* comprising supplying the cell with a double stranded RNA in an amount sufficient to specifically attenuate expression of the target gene, wherein one of the strands of the double stranded RNA is capable of hybridizing to the target gene *in vitro* in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 50°C, and provided that, when the double stranded RNA is supplied to the cell by delivery to the cell of double stranded RNA, the double stranded RNA is formed from single-stranded RNA that is purified in the absence of phenol or chloroform.
- 2. (original) The method of claim 1 wherein the target gene is an endogenous gene.
- 3. (original) The method of claim 1 wherein the target gene is a foreign gene.
- 4. (previously presented) The method of claim 1 wherein the target gene is a chromosomal gene.
- 5. (previously presented) The method of claim 1 wherein the target gene is an extrachromosomal gene.
- 6. (previously presented) The method of claim 1 wherein the target gene is from a pathogen capable of infecting the cell.
- 7. (original) The method of claim 6 wherein the pathogen is selected from the group consisting of a virus, bacterium, fungus or protozoan.
- 8. (canceled)

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9. (previously presented) The method of claim 1 wherein the vertebrate cell is a fish cell.

10. (previously presented) The method of claim 1 wherein the vertebrate cell is a mammalian

cell.

11-14. (canceled)

15. (previously presented) The method of claim 1 wherein the double stranded RNA

comprises a nucleotide sequence that is complementary to the nucleotide sequence of at least a

portion of the target gene.

16. (currently amended) The method of claim 1 wherein the double stranded RNA comprises

a nucleotide sequence that is complementary to a region of at least [50] about 25 bases of the

target gene.

17. (previously presented) The method of claim 1 wherein the double stranded RNA is

supplied in an amount sufficient to completely inhibit expression of the target gene.

18. (currently amended) The method of claim 1 in which the double stranded RNA

comprises a single strand which is comprising self-complementary portions.

19. (original) The method of claim 1 in which the double stranded RNA comprises two

separate complementary strands.

20. (currently amended) The method of claim 1 wherein the cell is an embryo embryonic

<u>cell</u>.

21. (currently amended) The method of claim 20 wherein the embryo embryonic cell is a fish

embryo embryonic cell.

- 22. (currently amended) The method of claim 20 wherein the embryo embryonic cell is supplied with the double stranded RNA using microinjection.
- 23. (currently amended) The method of claim 1 wherein the cell is present in a cell culture, a tissue, an organ, or an organism.
- 24. (previously presented) The method of claim 23 wherein the cell is present in an organism, and the cell is supplied with the double stranded RNA by introducing double stranded RNA into a body cavity or interstitial space of the organism.
- 25. (previously presented) The method of claim 23 wherein the cell is present in an organism, and wherein the cell is supplied with the double stranded RNA by delivering double stranded RNA to the organism via oral, topical, parenteral, vaginal, rectal, intranasal, ophthalmic, or intraperitoneal administration.
- 26. (canceled)
- 27. (previously presented) The method of claim 1 wherein supplying the double stranded RNA to the cell comprises delivering double-stranded RNA to the cell, and wherein the double stranded RNA is treated with RNAse prior to delivery to the cell.
- 28. (previously presented) The method of claim 1 wherein supplying the double stranded RNA to the cell comprises delivering double stranded RNA to the cell, the method further comprising, prior to delivering the double stranded RNA to the cell, annealing two complementary single stranded RNAs to yield the double stranded RNA.
- 29. (previously presented) The method of claim 28 wherein the single stranded RNAs are annealed in the presence of potassium chloride.

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30. (original) The method of claim 1 wherein the function of the target gene is unknown.

31. (previously presented) The method of claim 1 further comprising introducing into the cell

a second double stranded RNA in an amount sufficient to attenuate expression of a second target

gene, wherein one of the strands of the second double stranded RNA is capable of hybridizing to

the second target gene in vitro in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at

50°C.

32. (previously presented) The method of claim 1 comprising introducing into the cell

multiple double stranded RNAs in an amount sufficient to attenuate expression of multiple target

genes, wherein one strand of each double stranded RNA is capable of hybridizing to the

corresponding target gene in vitro in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at

50°C.

33-38. (canceled)

39. (currently amended) The method of claim 23 further comprising identifying a phenotypic

change in the <u>organism</u> cell culture associated with attenuated expression of the target gene.

40-47. (canceled)

48. (previously presented) A method for attenuating the expression of a target gene in a

vertebrate cell comprising:

annealing two complementary single stranded RNAs in the presence of potassium

chloride to yield double stranded RNA;

contacting the double stranded RNA with RNAse to purify the double stranded RNA by

removing single stranded RNA; and

introducing the purified double stranded RNA into the cell in an amount sufficient to

specifically attenuate expression of the target gene;

wherein one of the strands of the double stranded RNA is capable of hybridizing to the

target gene *in vitro* in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 50°C, and wherein the double stranded RNA is formed from single-stranded RNA that is purified in the absence of phenol or chloroform.

49-60. (canceled)

61. (withdrawn) A kit comprising reagents for attenuating the expression of a target gene in a cell, the kit comprising:

a DNA template comprising two different promoters selected from the group consisting of a T7 promoter, a T3 promoter and an SP6 promoter, each promoter operably linked to a nucleotide sequence, such that two complementary single stranded RNAs are capable of being transcribed from the DNA template;

a plurality of primers for amplification of the nucleotide sequence; nucleotide triphosphates for forming RNA;

at least two RNA polymerases, each capable of binding to a promoter on the DNA template and causing transcription of the nucleotide sequence to which the promoter is operably linked;

a purification column for purifying single stranded RNA;

buffer for annealing single stranded RNAs to yield double stranded RNA; and RNAse A or RNAse T for purifying double stranded RNA;

wherein one of the strands of the double stranded RNA is capable of hybridizing to the target gene *in vitro* in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 50°C.

- 62. (previously presented) The method of claim 1 wherein one of the strands of the double stranded RNA is capable of hybridizing to the target gene *in vitro* in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 70°C.
- 63. (currently amended) A method for attenuating the expression of a target gene in a vertebrate cell <u>in vivo</u> comprising delivering a double stranded RNA to the cell in an amount sufficient to specifically attenuate expression of the target gene, wherein the double stranded

RNA comprises a nucleotide sequence that is complementary to a region of at least 25 nucleotides of the target gene, and wherein the double stranded RNA is formed from single-stranded RNA that is purified in the absence of phenol or chloroform.

- 64. (withdrawn) A method for attenuating the expression of two or more target genes in a vertebrate cell, the method comprising supplying the cell with two or more double stranded RNAs in an amount sufficient to specifically attenuate expression of the target genes, wherein one strand of each double stranded RNA is capable of hybridizing to the corresponding target gene *in vitro* in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 50°C.
- 65. (withdrawn) A method for attenuating the expression of two or more target genes in a vertebrate cell, the method comprising supplying the cell with two or more double stranded RNAs in an amount sufficient to specifically attenuate expression of the target genes, wherein one strand of each double stranded RNA comprises a nucleotide sequence that is complementary to a region of at least 25 nucleotides of a target gene.
- 66. (withdrawn) A method for reducing or preventing an immune response in a recipient organism to a transplant tissue obtained from a donor organism, the method comprising supplying the transplant tissue with a double stranded RNA *in vitro* prior to implanting the transplant tissue into the recipient organism, wherein the double stranded RNA attenuates the expression of a target gene in the transplant tissue that can elicit an immune response in a recipient, and wherein one of the strands of the double stranded RNA is capable of hybridizing to the target gene *in vitro* in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 50°C.
- 67. (withdrawn) A pharmaceutical composition for inhibiting the function of a target gene in a vertebrate cell, wherein the composition comprises a double stranded RNA in an amount sufficient to specifically attenuate expression of the target gene, wherein the double stranded RNA comprises a nucleotide sequence that is complementary to a region of at least 25 nucleotides of the target gene, and wherein the composition does not comprise phenol or chloroform.

- 68. (previously presented) The method of any one of claims 48, 63, 64, 65 or 66 wherein the cell is a mammalian cell.
- 69. (previously presented) The method of any one of claims 1, 48, 63, 64, 65 or 66 wherein the cell is a human cell.
- 70. (previously presented) The method of any one of claims 23, 48, 63, 64, 65 or 66 wherein the cell is present in a mammal.
- 71. (previously presented) The method of claim 70 wherein the cell is present in a human.
- 72. (previously presented) The method of claim 63 or 65 wherein the target gene is associated with a disease.
- 73. (previously presented) The method of claim 63 or 65 wherein the target gene is associated with a disease from a pathogen.
- 74. (previously presented) The method of claim 1, 64, 65 or 66 wherein the double stranded RNA is supplied to the cell by delivering to the cell a DNA encoding the double stranded RNA.
- 75. (new) A method for attenuating the expression of a target gene in a vertebrate cell *ex vivo* comprising:

explanting a cell from a vertebrate organism;

supplying the cell with a double stranded RNA in an amount sufficient to specifically attenuate expression of the target gene, wherein one of the strands of the double stranded RNA is capable of hybridizing to the target gene in vitro in 400 mM NaCl, 40 mM PIPES pH 6.4, and 1 mM EDTA, at 50°C, and provided that, when the double stranded RNA is supplied to the cell by delivery to the cell of double stranded RNA, the double stranded RNA is formed from single-stranded RNA that is purified in the absence of phenol or chloroform; and

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implanting the cell into a vertebrate organism.

76. (new) The method of claim 75, wherein the cell is implanted back into the vertebrate organism from which it was explanted.

- 77. (new) The method of claim 1, wherein the double stranded RNA comprises overhanging ends.
- 78. (new) The method of claim 1, wherein the double stranded RNA has a length of less than about 200 bases.
- 79. (new) The method of claim 18 wherein the double stranded RNA comprises a nucleotide sequence that is complementary to a region of at least about 25 bases of the target gene.
- 80. (new) The method of claim 19 wherein the double stranded RNA comprises a nucleotide sequence that is complementary to a region of at least about 25 bases of the target gene